SEQ ID NO: 1

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  1 GCAGCCCTGC CTCAGAAAAC AGAAGGACGC AGCACACTCA CGGTGACTCA CCCCCATGTG   60
 61 GCTGGGAGGNG AGGGAGCCTC CTGAGGCCAGG GCCAGGGCCAG CCGTCAGGTG GGTGACGGCA  120
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121 GGGGTCTTGC CATGGTGGGC ACAGGGGCTG CATACAGCTT ACTCAGTGAC AATCGAGTCC  180
181 CTGGTGCCCAG CCTCTGGAAG TCTGGAAGTG AGCAATGTTT CCCATTAAGG AAAGTGTGTG  240
241 GCCGGCCATG CCCCCCAACG TTGCACACTC ACTGCCTTTG CAGGGTTGGG GCTTCCAGTC  300
301 ACAGGGTCCC ATCCACGTAC CAGCCCAGGT GGCTGCAGAA GGTCCCTCGC AGTCATGAAA  360
361 CCAAGGGAGG CTTGGGAAAC CACATCTGAA GGGCATGGCT TTGATTTAGT GAGAGGGTGG  420
421 GGCTGGGCTG GGCAAGGCCA CCAGGTCTGA GTCAGAGCCA GAGGCAGGAA GCTGGTCCCC  480
481 AGCACTGCCCC GCCGCCTCTG CGATGCAGTC CTCCTGGCCA CCTGAGAACA GCCTGTAGAG  540
541 AGGCAGTGGC GTCTTTCGGA CTTC                                          564
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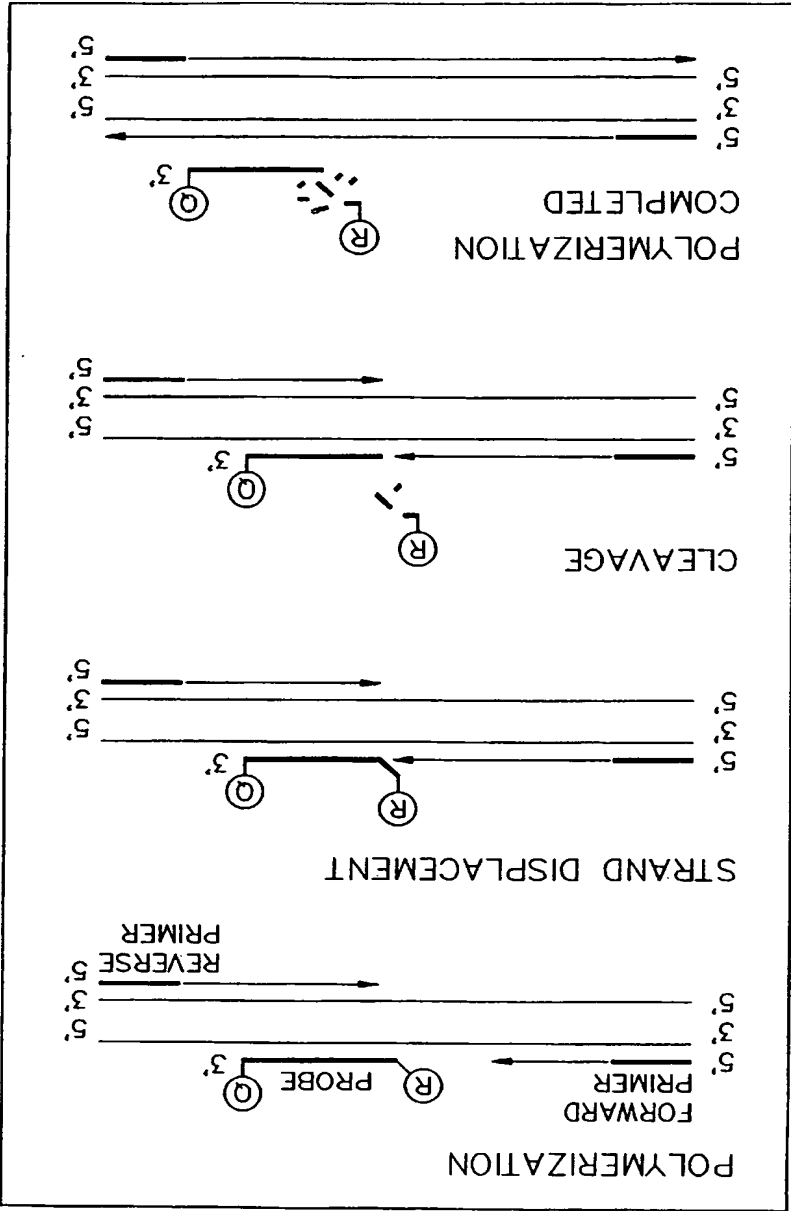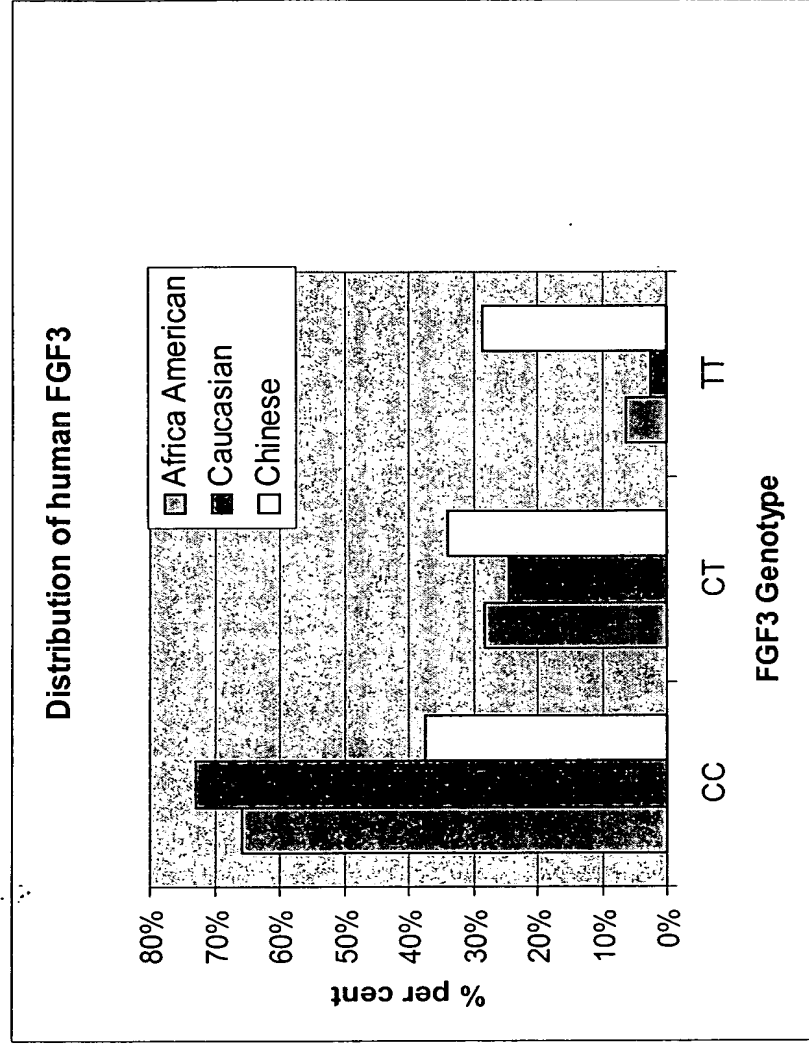
* C-allele: N=C
  T-allele: N=T

Figure 1

Figure 2

Distribution of human FGF3

% per cent

FGF3 Genotype

Legend:
- Africa American
- Caucasian
- Chinese

Figure 3